US-08-864-038A-3

Query Match 65.9%; Score 170; DB 3; Length 738;
Best Local Similarity 70.2%; Pred. No. 3.7e-08;
Matches 33; Conservative 1; Mismatches 11; Indels 2; Gaps 1;

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Qy       1 SGGSGSGGQSGGSGSGGQSGGSGSGGQSGGSGSGGQSGGSGSGGQSG 47
           :|| ||||  |||| ||  |||| || ||||| || ||||| ||  |
Db     665 NGGGGSGG--GGSGGGGSGGGSGGGGGSGGSGGGGGSGGSGGGGSGG 709
```

RESULT 2
US-08-529-190B-1
; Sequence 1, Application US/08529190B
; Patent No. 5833991
; GENERAL INFORMATION:
; APPLICANT: Masucci, Maria G.
; TITLE OF INVENTION: GLYCINE-CONTAINING SEQUENCES
; TITLE OF INVENTION: CONFERRING INVISIBILITY TO THE IMMUNE SYSTEM
; NUMBER OF SEQUENCES: 76
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Banner & Witcoff, Ltd.
; STREET: One Financial Center
; CITY: Boston
; STATE: MA
; COUNTRY: USA
; ZIP: 02111
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Diskette
; COMPUTER: IBM Compatible
; OPERATING SYSTEM: DOS
; SOFTWARE: Wordperfect 6.1
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/529,190B
; FILING DATE: 15-SEP-1995
; CLASSIFICATION: 514
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: SE9501324-9
; FILING DATE: 10-APR-1995
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: US08/522,595
; FILING DATE: 01-SEP-1995
; ATTORNEY/AGENT INFORMATION:
; NAME: Williams, Ph.D., Kathleen A
; REGISTRATION NUMBER: 34,380
; REFERENCE/DOCKET NUMBER: 3255/53015
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 617-345-9100
; TELEFAX: 617-345-9111
; INFORMATION FOR SEQ ID NO: 1:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 235 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: unknown
; MOLECULE TYPE: protein
US-08-529-190B-1

Query Match 59.7%; Score 154; DB 2; Length 235;
Best Local Similarity 54.3%; Pred. No. 3.5e-07;
Matches 25; Conservative 13; Mismatches 8; Indels 0; Gaps 0;

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Qy       2 GGSGSGGQSGGSGSGGQSGGSGSGGQSGGSGSGGQSGGSGSGGQSG 47
             ||:|:|| :||:|:|| :||:|  |  ||:|:|| :||:|:|| :|
Db      57 GGAGAGGGAGGAGAGGGAGGAGGAGAGGGAGAGGGAGGAGAGGGAG 102
```

RESULT 3
US-09-249-585A-3
; Sequence 3, Application US/09249585A
; Patent No. 6417002
; GENERAL INFORMATION:
; APPLICANT: Horlick, Robert
; TITLE OF INVENTION: METHOD FOR MAINTENANCE AND SELECTION OF EPISOMES
; FILE REFERENCE: 0867/0D905
; CURRENT APPLICATION NUMBER: US/09/249,585A
; CURRENT FILING DATE: 1999-02-11
; NUMBER OF SEQ ID NOS: 18
; SOFTWARE: PatentIn version 3.0
; SEQ ID NO 3
; LENGTH: 641
; TYPE: PRT
; ORGANISM: Epstein Barr Virus
US-09-249-585A-3

Query Match 59.7%; Score 154; DB 4; Length 641;
Best Local Similarity 54.3%; Pred. No. 8e-07;
Matches 25; Conservative 13; Mismatches 8; Indels 0; Gaps 0;

```
Qy       2 GGSGSGGQSGGSGSGGQSGGSGSGGQSGGSGSGGQSGGSGSGGQSG 47
             ||:|:|| :||:|:|| :||:|  |  ||:|:|| :||:|:|| :|
Db     146 GGAGAGGGAGGAGAGGGAGGAGGAGAGGGAGAGGGAGGAGAGGGAG 191
```

RESULT 4
US-09-052-995-1
; Sequence 1, Application US/09052995
; Patent No. 6183956
; GENERAL INFORMATION:
; APPLICANT: Sivaraja, Mohanram
; APPLICANT: Strulovici, Berta
; APPLICANT: Flores, Osvaldo A.
; TITLE OF INVENTION: High Throughput In Vitro Screening Assay
; TITLE OF INVENTION: for Transcription Modulators
; NUMBER OF SEQUENCES: 13
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Townsend and Townsend and Crew LLP
; STREET: Two Embarcadero Center, Eighth Floor
; CITY: San Francisco
; STATE: California
; COUNTRY: USA
; ZIP: 94111-3834
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.30
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/09/052,995
; FILING DATE: 31-MAR-1998
; CLASSIFICATION: 435
; ATTORNEY/AGENT INFORMATION:
; NAME: Parent, Annette S.
; REGISTRATION NUMBER: 42,058
; REFERENCE/DOCKET NUMBER: 018781-000600US
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (415) 576-0200
; TELEFAX: (415) 576-0300
; INFORMATION FOR SEQ ID NO: 1:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 201 amino acids
; TYPE: amino acid
; STRANDEDNESS:
; TOPOLOGY: linear
; MOLECULE TYPE: peptide
US-09-052-995-1

Query Match 57.0%; Score 147; DB 4; Length 201;
Best Local Similarity 56.5%; Pred. No. 1.2e-06;
Matches 26; Conservative 0; Mismatches 20; Indels 0; Gaps 0;

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Qy       2 GGSGSGGQSGGSGSGGQSGGSGSGGQSGGSGSGGQSGGSGSGGQSG 47
             || | ||  || | ||  || | ||  || | ||  || | ||  |
Db      58 GGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGPGG 103
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